

GenCore version 6.3  
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:09:48 ; Search time 1274 Seconds  
 (without alignments)  
 13572.847 Million cell

updates/sec

Title: US-09-300-482-311  
 Perfect score: 274  
 Sequence: 1  
 cttacattcctttctccacc.....gccaaagcatgccgtcgaccg 274

Scoring table: IDENTITY\_NUC  
 Gapop 10.0 , Gapext 1.0

Searched: 111716937 seqs, 31560637849 residues

Total number of hits satisfying chosen parameters: 223433874

Minimum DB seq length: 0  
 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
 Maximum Match 100%  
 Listing first 45 summaries

Database : Pending\_Patents\_NA\_Main:\*  
 1: /ABSS/Data/CRF/ptodata/1/pna/PCTUSA\_COMB.seq:\*  
 2: /ABSS/Data/CRF/ptodata/1/pna/PCTUSB\_COMB.seq:\*  
 3: /ABSS/Data/CRF/ptodata/1/pna/PCTUSC\_COMB.seq:\*  
 4: /ABSS/Data/CRF/ptodata/1/pna/PCTUSD\_COMB.seq:\*  
 5: /ABSS/Data/CRF/ptodata/1/pna/US075\_COMB.seq:\*  
 6: /ABSS/Data/CRF/ptodata/1/pna/US076\_COMB.seq:\*  
 7: /ABSS/Data/CRF/ptodata/1/pna/US077\_COMB.seq:\*  
 8: /ABSS/Data/CRF/ptodata/1/pna/US078\_COMB.seq:\*  
 9: /ABSS/Data/CRF/ptodata/1/pna/US079\_COMB.seq:\*  
 10: /ABSS/Data/CRF/ptodata/1/pna/US080\_COMB.seq:\*  
 11: /ABSS/Data/CRF/ptodata/1/pna/US081\_COMB.seq:\*  
 12: /ABSS/Data/CRF/ptodata/1/pna/US082\_COMB.seq:\*  
 13: /ABSS/Data/CRF/ptodata/1/pna/US083\_COMB.seq:\*  
 14: /ABSS/Data/CRF/ptodata/1/pna/US084\_COMB.seq:\*  
 15: /ABSS/Data/CRF/ptodata/1/pna/US085\_COMB.seq:\*